-continued

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Gly Ser Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp Lys Ser Arg Trp
                405                 410                 415

Gln Glu Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His
            420                 425                 430

Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Leu Gly Lys
        435                 440                 445
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1. An antigen-binding molecule, optionally isolated, which is capable of binding to VISTA and inhibiting VISTA-mediated signalling, independently of Fc-mediated function.

2. The antigen-binding molecule according to claim **1**, wherein the antigen-binding molecule comprises:

(i) a heavy chain variable (VH) region incorporating the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:305

HC-CDR2 having the amino acid sequence of SEQ ID NO:306

HC-CDR3 having the amino acid sequence of SEQ ID NO:307; and

(ii) a light chain variable (VL) region incorporating the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:308

LC-CDR3 having the amino acid sequence of SEQ ID NO:43.

3. The antigen-binding molecule according to claim **1** or claim **2**, wherein the antigen-binding molecule comprises:

(i) a heavy chain variable (VH) region incorporating the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:290

HC-CDR2 having the amino acid sequence of SEQ ID NO:291

HC-CDR3 having the amino acid sequence of SEQ ID NO:278; and

(ii) a light chain variable (VL) region incorporating the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:309

LC-CDR3 having the amino acid sequence of SEQ ID NO:43.

4. The antigen-binding molecule according to any one of claims **1** to **3**, wherein the antigen-binding molecule comprises:

(i) a heavy chain variable (VH) region incorporating the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:290

HC-CDR2 having the amino acid sequence of SEQ ID NO:291

HC-CDR3 having the amino acid sequence of SEQ ID NO:278; and

(ii) a light chain variable (VL) region incorporating the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:295

LC-CDR3 having the amino acid sequence of SEQ ID NO:43.

5. The antigen-binding molecule according to any one of claims **1** to **3**, wherein the antigen-binding molecule comprises:

(i) a heavy chain variable (VH) region incorporating the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:290

HC-CDR2 having the amino acid sequence of SEQ ID NO:291

HC-CDR3 having the amino acid sequence of SEQ ID NO:278; and

(ii) a light chain variable (VL) region incorporating the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:300

LC-CDR3 having the amino acid sequence of SEQ ID NO:43.

6. The antigen-binding molecule according to any one of claims **1** to **5**, wherein the antigen-binding molecule comprises:

a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and

a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:310.

7. The antigen-binding molecule according to any one of claims **1** to **6**, wherein the antigen-binding molecule comprises:

a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and

a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:294.

8. The antigen-binding molecule according to any one of claims **1** to **6**, wherein the antigen-binding molecule comprises:

a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and